
Sequence Listing was accepted.

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Timestamp: [year=2008; month=9; day=19; hr=16; min=11; sec=22; ms=759;]

Validated By CRFValidator v 1.0.3

Application No: 10540959 Version No: 2.0

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Finished: 2008-08-21 20:25:41.003

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Actual SeqID Count: 9

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Arg Gly Ser Cys Ser Leu Phe Thr Cys Gln Asn Gly Ile Val Trp Thr
                       55
Asn Gly Thr His Val Thr Tyr Arg Lys Asp Thr Arg Tyr Lys Leu Leu
                   70
                                       75
Gly Asp Leu Ser Arg Arg Asp Val Ser Leu Thr Ile Glu Asn Thr Ala
                                   90
Val Ser Asp Ser Gly Val Tyr Cys Cys Arg Val Glu His Arg Gly Trp
                              105
          100
Phe Asn Asp Met Lys Ile Thr Val Ser Leu Glu Ile Val Pro Pro Lys
Val Thr Thr Thr Pro Ile Val Thr Thr Val Pro Thr Val Thr Thr Val
                       135
                                           140
Arg Thr Ser Thr Thr Val Pro Thr Thr Thr Thr Val Pro Thr Thr
                   150
                                       155
Val Pro Thr Thr Met Ser Ile Pro Thr Thr Thr Thr Val Pro Thr Thr
                                   170
Met Thr Val Ser Thr Thr Ser Val Pro Thr Thr Ser Ile Pro
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                              185
Thr Thr Ser Val Pro Val Thr Thr Val Ser Thr Phe Val Pro
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Pro Met Pro Leu Pro Arg Gln Asn His Glu Pro Val Ala Thr Ser Pro

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Ile A	Arg	Arg	Glu	Pro	Thr	Ser	Ser	Pro	Leu 250	Tyr	Ser	Tyr	Thr	Thr 255	Asp
Gly A	Asn	Asp			Thr	Glu	Ser			Gly	Leu	Trp			Asn
	_,		260	_,	_			265	_	_	_,		270	_,	
Gln T	inr	275	Leu	Pne	Leu	Glu	280	ser	Leu	Leu	Thr	285	Asn	Thr	Thr
Lys G	31y 290	Ile	Tyr	Ala	Gly	Val 295	Cys	Ile	Ser	Val	Leu 300	Val	Leu	Leu	Ala
Leu I 305	Leu	Gly	Val	Ile	Ile 310	Ala	Lys	Lys	Tyr	Phe 315	Phe	Lys	Lys	Glu	Val 320
Gln G	Gln	Leu	Ser	Val 325		Phe	Ser	Ser	Leu 330		Ile	Lys	Ala	Leu 335	
Asn A	Ala	Val			Glu	Val	Gln			Asp	Asn	Ile			Glu
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Ser V	/al	Ala	Gly 20	Ser	Val	Lys	Val	Gly 25	Gly	Glu	Ala	Gly	Pro 30	Ser	Val
Thr I	Leu	Pro 35		His	Tyr	Ser	Gly		Val	Thr	Ser	Met 45		Trp	Asn
Arg G	-		Cys	Ser	Leu			Cys	Gln	Asn	_		Val	Trp	Thr
Asn G	50 Gly	Thr	His	Val	Thr	55 Tyr	Arg	Lys	Asp	Thr	60 Arg	Tyr	Lys	Leu	Leu
65 Gly A	Asp	Leu	Ser	Arg	70 Arg	Asp	Val	Ser	Leu	75 Thr	Ile	Glu	Asn	Thr	80 Ala
				85		_			90					95	
Val S	Ser	Asp	Ser 100	Gly	Val	Tyr	Cys	Cys 105	Arg	Val	Glu	His	Arg 110	Gly	Trp
Phe A	Asn	Asp	Met	Lys	Ile	Thr	Val 120	Ser	Leu	Glu	Ile	Val 125	Pro	Pro	Lys
Val T	Chr L30	Thr	Thr	Pro	Ile	Val	Thr	Thr	Val	Pro	Thr	Val	Thr	Thr	Val
Arg I		Ser	Thr	Thr			Thr	Thr	Thr			Pro	Thr	Thr	
145 Val E	Pro	Thr	Thr	M≏+	150 Ser	T1_	Pro	Thr	Thr	155 Thr	Thr	Val	Pro	Thr	160 Thr
				165					170					175	
Met I	Γhr	Val	Ser 180	Thr	Thr	Thr	Ser	Val 185	Pro	Thr	Thr	Thr	Ser 190	Ile	Pro
Thr I	Γhr		Ser	Val	Pro	Val		Thr	Thr	Val	Ser		Phe	Val	Pro
Dr	1 0 ±	195	т с	D ====	7	C1	200	U	C1	D	77-7	205	Th	C	Dana
	210				-	215					220				
Ser S	Ser	Pro	Gln	Pro	Ala	Glu	Thr	His	Pro	Thr	Thr	Leu	Gln	Gly	Ala

225 230 230 235 240 240 Ile Arg Arg Glu Pro Thr Ser Ser Pro Leu Tyr Ser Tyr Thr Thr Asp

245 250 255

Gly Asn Asp Thr Val Thr Glu Ser Ser Asp Gly Leu Trp Asn Asn Asn 260 265 270

Gln Thr Gln Leu Phe Leu Glu His Ser Leu Leu Thr Ala Asn Thr Thr 275 280 285

Lys Gly Ile Tyr Ala Gly Val Cys Ile Ser Val Leu Val Leu Leu Ala 290 295 300

Leu Leu Gly Val Ile Ile Ala Lys Lys Tyr Phe Phe Lys Lys Glu Val 305 310 315 320

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Thr Leu Pro Cys His Tyr Ser Gly Ala Val Thr Ser Met Cys Trp Asn 35 40 45

Arg Gly Ser Cys Ser Leu Phe Thr Cys Gln Asn Gly Ile Val Trp Thr 50 55 60

Asn Gly Thr His Val Thr Tyr Arg Lys Asp Thr Arg Tyr Lys Leu Leu 65 70 75 80

Gly Asp Leu Ser Arg Asp Val Ser Leu Thr Ile Glu Asn Thr Ala 85 90 95

Val Ser Asp Ser Gly Val Tyr Cys Cys Arg Val Glu His Arg Gly Trp
100 105 110

Phe Asn Asp Met Lys Ile Thr Val Ser Leu Glu Ile Val Pro Pro Lys 115 120 125

Val Thr Thr Thr Pro Ile Val Thr Thr Val Pro Thr Val Thr Thr Val 130 135 140

Arg Thr Ser Thr Thr Val Pro Thr Thr Thr Thr Val Pro Thr Thr 145 150 155 160

Val Pro Thr Thr Met Ser Ile Pro Thr Thr Thr Thr Val Pro Thr Thr

165 170 175

Met Thr Val Ser Thr Thr Thr Ser Val Pro Thr Thr Thr Ser Ile Pro 180 185 190

Thr Thr Thr Ser Val Pro Val Thr Thr Thr Val Ser Thr Phe Val Pro 195 200 205

Pro Met Pro Leu Pro Arg Gln Asn His Glu Pro Val Ala Thr Ser Pro 210 215 220

Ser Ser Pro Gln Pro Ala Glu Thr His Pro Thr Thr Leu Gln Gly Ala 225 230 235 240

Ile Arg Arg Glu Pro Thr Ser Ser Pro Leu Tyr Ser Tyr Thr Thr Asp
245 250 255

Gly Asn Asp Thr Val Thr Glu Ser Ser Asp Gly Leu Trp Asn Asn Asn 260 265 270

Gln Thr Gln Leu Phe Leu Glu His Ser Leu Leu Thr Ala Asn Thr Thr

275 280 285

Lys	Gly 290	Val	Asp	Lys	Thr	His 295	Thr	Cys	Pro	Pro	Cys	Pro	Ala	Pro	Glu
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Val	Ser	His	Glu	325 Asp	Pro	Glu	Val	Lys	330 Phe	Asn	Trp	Tyr	Val	335 Asp	Gly
			340					345					350		
Val	Glu	Val 355	His	Asn	Ala	Lys	Thr 360	Lys	Pro	Arg	Glu	Glu 365	Gln	Tyr	Asn
Ser	Thr 370	Tyr	Arg	Val	Val	Ser 375	Val	Leu	Thr	Val	Leu 380	His	Gln	Asp	Trp
Leu 385	Asn	Gly	Làs	Glu	Tyr 390	Lys	Cys	Lys	Val	Ser	Asn	Lys	Ala	Leu	Pro 400
	Pro	Ile	Glu	_		Ile	Ser	Lys			Gly	Gln	Pro		
Pro	Gln	Val	Tur	405 Thr	T.e11	Pro	Pro	Ser	410 Ara	Asp	Glu	T.e11	Thr	415	Asn
			420					425	_	_			430	_	
Gln	Val	Ser	Leu	Thr	Cys	Leu		Lys	Gly	Phe	Tyr		Ser	Asp	Ile
Ala	Val		Trp	Glu	Ser	Asn	440 Glv	Gln	Pro	Glu	Asn	445 Asn	Tyr	Lvs	Thr
1120	450	0_0	ттр	0_0	501	455	019	0211		0_4	460	11511	- 1 -	2,0	
Thr	Pro	Pro	Val	Leu	Asp	Ser	Asp	Gly	Ser	Phe	Phe	Leu	Tyr	Ser	Lys
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Leu	Thr	Val	Asp	_	Ser	Arg	Trp	Gln		Gly	Asn	Val	Phe		Cys
_				485		_			490	_				495	_
Ser	Val	Met	H1S	GLu	Ala	Leu	His	Asn 505	HIS	Tyr	Thr	GIn	Lуs 510	Ser	Leu
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Thr	Leu	Pro 35	CÀR	His	Tyr	Ser	Gly 40	Ala	Val	Thr	Ser	Met 45	Суз	Trp	Asn
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_	_			85	-	_			90					95	
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Phe Asn Asp Met Lys Ile Thr Val Ser Leu Glu Ile Val Pro Pro Lys

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Leu Gly Gly Pro Ser Val Phe Leu Phe Pro Pro Lys Pro Lys Asp Thr
                150
                             155
Leu Met Ile Ser Arg Thr Pro Glu Val Thr Cys Val Val Val Asp Val
              165
                                 170
Ser His Glu Asp Pro Glu Val Lys Phe Asn Trp Tyr Val Asp Gly Val
           180 185
Glu Val His Asn Ala Lys Thr Lys Pro Arg Glu Glu Gln Tyr Asn Ser
                          200
Thr Tyr Arg Val Val Ser Val Leu Thr Val Leu His Gln Asp Trp Leu
                      215
Asn Gly Lys Glu Tyr Lys Cys Lys Val Ser Asn Lys Ala Leu Pro Ala
                 230
                                    235
Pro Ile Glu Lys Thr Ile Ser Lys Ala Lys Gly Gln Pro Arq Glu Pro
                                 250
              245
Gln Val Tyr Thr Leu Pro Pro Ser Arg Asp Glu Leu Thr Lys Asn Gln
                             265
                                                270
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Val Ser Leu Thr Cys Leu Val Lys Gly Phe Tyr Pro Ser Asp Ile Ala
                          280
Val Glu Trp Glu Ser Asn Gly Gln Pro Glu Asn Asn Tyr Lys Thr Thr
                      295
                                         300
Pro Pro Val Leu Asp Ser Asp Gly Ser Phe Phe Leu Tyr Ser Lys Leu
        310
                            315
Thr Val Asp Lys Ser Arg Trp Gln Gln Gly Asn Val Phe Ser Cys Ser
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Val Thr Thr Thr Pro Ile Val Thr Thr Val Pro Thr Val Thr Thr Val 135 140 Arg Thr Ser Thr Thr Val Pro Thr Thr Thr Thr Val Pro Thr Thr 145 150 155

Val Pro Thr Thr Met Ser Ile Pro Thr Thr Thr Thr Val Pro Thr Thr 165 170

Met Thr Val Ser Thr Thr Thr Ser Val Pro Thr Thr Ser Ile Pro 185

Thr Thr Ser Val Pro Val Thr Thr Val Ser Thr Phe Val Pro 200

Pro Met Pro Leu Pro Arg Gln Asn His Glu Pro Val Ala Thr Ser Pro 215

Ser Ser Pro Gln Pro Ala Glu Thr His Pro Thr Thr Leu Gln Gly Ala 230 235

Ile Arg Arg Glu Pro Thr Ser Ser Pro Leu Tyr Ser Tyr Thr Thr Asp 245 250

Gly Asn Asp Thr Val Thr Glu Ser Ser Asp Gly Leu Trp Asn Asn Asn 265

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Lys Gly Val Glu His His His His His 295

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                                                                      180
caaaatacac ttatttggac caatggacat cgtgtcacct atcagaagag cagtcggtac
                                                                       240
aacttaaagg ggcatatttc agaaggagat gtgtccttga cgatagagaa ctctgttgag
                                                                      300
agtgacagtg gtctgtattg ttgtcgagtg gagattcctg gatggtttaa tgatcagaaa
                                                                       360
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                                                                      480
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                                                                      600
tcccatactc ctacagactg gaatggcact gcgacatcct caggagatac ctggagtaat
                                                                       660
                                                                      720
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                                                                      900
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